(FILE 'HOME' ENTERED AT 14:17:03 ON 26 APR 2006)

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FILE 'MEDLINE, AGRICOLA, BIOSIS' ENTERED AT 14:17:18 ON 26 APR 2006
        3115391 S PLANT
L1
                                                                            1541.10G
H240G
        3474980 S DNA OR GENE OR POLYNUCL?
L2
        1226032 S TRANSFOR? OR TRANSFECT? OR RECOMBIN? OR BIOLISTIC OR AGROBACT
L3
L4
         279511 S PROMOTER
L5
L6
             13 S L4 AND L3
              7 DUP REM L6 (6 DUPLICATES REMOVED)
L7
=> type 17 1-7 all
                       MEDLINE on STN
     ANSWER 1 OF 7
L7
     2001461223
                    MEDLINE
AN
DN
     PubMed ID: 11506359
TI
    Glucocorticoid-inducible gene expression in rice.
     Ouwerkerk P B; de Kam R J; Hoge J H; Meijer A H
ΑU
     Institute of Molecular Plant Sciences, Leiden University, Clusius
     Laboratory, The Netherlands.
     Planta, (2001 Jul) Vol. 213, No. 3, pp. 370-8.
SO
     Journal code: 1250576. ISSN: 0032-0935.
CY
     Germany: Germany, Federal Republic of
     Journal; Article; (JOURNAL ARTICLE)
DT
LΑ
     English
FS
     Priority Journals
     GENBANK-AF294979; GENBANK-AF294980; GENBANK-AF294981; GENBANK-AF294982
OS
EΜ
     200206
    Entered STN: 20 Aug 2001
     Last Updated on STN: 25 Jun 2002
     Entered Medline: 24 Jun 2002
     We have studied the use of a glucocorticoid receptor-based inducible gene
AΒ
     expression system in the monocotyledonous model plant rice (Oryza sativa
     L.). This system, originally developed by T. Aoyama and N.-H. Chua
     [(1997) Plant J 11: 605-612], is based on the chimaeric transcriptional
     activator GVG, consisting of the yeast Gal4 DNA-binding domain, the VP16
     activation domain and the glucocorticoid receptor domain. For application
     in rice, we designed an optimized binary vector series (pINDEX) and tested
     this with the beta-glucuronidase (gusA) reporter gene. GUS expression was
     tightly controlled and relatively low concentrations (1-10 microM) of the
     glucocorticoid hormone dexamethasone (DEX) were able to induce GUS
     activities to levels comparable to those conferred by the strong
     cauliflower mosaic virus (CaMV) 35S promoter. DEX was taken up
     efficiently by the roots of tissue-cultured plantlets or mature plants in
     hydroponic culture, and induced GUS activity throughout the whole plant.
     DEX-induced GUS expression patterns were consistent in all lines and their
     T1 progeny. The phenotype of tissue-cultured rice plantlets was not
     affected when inductions with 10-100 microM DEX were limited to 1-4 days
     or when 2-week inductions were performed with 1 microM DEX, which was
     already sufficient to reach near-maximal GUS activity. However, 2-week
     inductions with 10 microM DEX caused growth retardation and developmental
     defects. As the severity of these effects varied between different lines,
     we could select lines with a mild phenotype for future use as activator
     lines in crosses with 'target' plants.
CT
     Base Sequence
     *Cloning, Molecular: MT, methods
      DNA-Binding Proteins: BI, biosynthesis
     *Dexamethasone: PD, pharmacology
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WEST Search History

Hide Items	Restore	Clear	Cancel

DATE: Wednesday, April 26, 2006

DATE: Wednesday, April 20, 2000				.<
Hide?	Set Name DB=PGPB	Query ,USPT,DWPI; PLUR=	Hit Count YES; OP=OR	10/541,315
	L9 .	L8 or 17	14 ·	1010
	L8	inze-dirk.in.	13	109
	L7	hatzfeld-yves.in.	3	1/2/61
	L6	11 adj3 13	24	W N
	L5	(11 or 12) adj5 13	1046	\mathcal{P}^{Q}
	L4	(11 or 12) and 13	14519	•
	L3	promoter	160705	·
П	L2	rice	127198	
	Ll	gos2	101	

END OF SEARCH HISTORY